

T. reesei eg18 nucleotide sequence (coding and non-coding)

[illegible]

MRATSLLLAAA LAVAGDALAG KIKYLGAIP GIDFGCDIDG SCPTDTSVP LLSYKGDGA GQMKHFAEDD
 10 20 30 40 50 60 70
 GLNVFRISAT WQFVLNNTVD GKLDLWGS YNKVYNACLE TGA YCMIDMH NFARYNGGII GQGGVSDDIF
 80 90 100 110 120 130 140
 VDLWVQIAKY YEDNDKIIFG LMNEPHDIDI EIWAQTCQV VTAIRKAGAT SQMILLPGTN FASVETVYST
 150 160 170 180 190 200 210
 GSAEALGKIT NPDGSTDLLY FDVHKYLDIN NSGSHAECTT DNVDADFNDFA DWLRONKROA IISSETGASME
 220 230 240 250 260 270 280
 PSCWTAFCQA NKAISENSDV YIGFVGWGAG SFDTSYILTL TPLGKPGNYT DNKLMNECIL DQFTLDEKYR
 290 300 310 320 330 340 350
 PTPTSISTAA BETATATATS DGDAPSTTKP IFRETASPT PNAVTKPSPD TSDSSDDDKD SAAASMSAQL
 360 370 380 390 400 410 420
 TGTVLFTVAA LGYMLVAF
 430 438

Figure 2
Predicted T. reesei EG VIII amino acid sequence